

Resource Summary Report

Generated by [RRID](#) on Aug 12, 2026

PermutMatrix

RRID:SCR_026979

Type: Tool

Proper Citation

PermutMatrix (RRID:SCR_026979)

Resource Information

URL: <http://www.atgc-montpellier.fr/permutmatrix/>

Proper Citation: PermutMatrix (RRID:SCR_026979)

Description: Software package for analysing and visualising data. Graphical environment to arrange gene expression profiles in optimal linear order.

Resource Type: software toolkit, software resource

Defining Citation: [PMID:15546938](#)

Keywords: Graphical environment, arrange gene expression profiles, optimal linear order,

Availability: Free, Available for download, Freely available

Resource Name: PermutMatrix

Resource ID: SCR_026979

Record Creation Time: 20250522T060228+0000

Record Last Update: 20260810T040935+0000

Ratings and Alerts

No rating or validation information has been found for PermutMatrix.

No alerts have been found for PermutMatrix.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 5 mentions in open access literature.

Listed below are recent publications. The full list is available at [RRID](#) .

Halo BA, et al. (2025) Drought-induced microbial dynamics in cowpea rhizosphere: Exploring bacterial diversity and bioinoculant prospects. PloS one, 20(3), e0320197.

Shin MK, et al. (2024) Nonsense-mediated mRNA decay of mRNAs encoding a signal peptide occurs primarily after mRNA targeting to the endoplasmic reticulum. Molecules and cells, 47(4), 100049.

Vily-Petit J, et al. (2022) Intestinal gluconeogenesis shapes gut microbiota, fecal and urine metabolome in mice with gastric bypass surgery. Scientific reports, 12(1), 1415.

He Q, et al. (2020) Dynamic Changes of the Anthocyanin Biosynthesis Mechanism During the Development of Heading Chinese Cabbage (*Brassica rapa* L.) and Arabidopsis Under the Control of BrMYB2. Frontiers in plant science, 11, 593766.

Nuthikattu S, et al. (2019) The Western Diet Regulates Hippocampal Microvascular Gene Expression: An Integrated Genomic Analyses in Female Mice. Scientific reports, 9(1), 19058.